

L. Lee

1645

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RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/151,409A

DATE: 04/15/1999  
TIME: 14:32:14

Input Set: I151409A.RAW

This Raw Listing contains the General Information  
Section and up to first 5 pages.

1 <110> APPLICANT: Dale, James B.  
2 <120> TITLE OF INVENTION: GROUP A STREPTOCOCCAL VACCINES  
3 <130> FILE REFERENCE: 481112.410  
4 <140> CURRENT APPLICATION NUMBER: US/09/151,409A  
5 <141> CURRENT FILING DATE: 1998-09-10  
6 <150> EARLIER APPLICATION NUMBER: US 60/058,635  
7 <151> EARLIER FILING DATE: 1997-09-12  
8 <160> NUMBER OF SEQ ID NOS: 16  
9 <170> SOFTWARE: PatentIn Ver. 2.0  
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165 1 5 10 15
166 caa gaa cgt gct gac aag ttt gag ata gaa aac aat acg tta aaa ctt 96
167 Gln Glu Arg Ala Asp Lys Phe Glu Ile Glu Asn Asn Thr Leu Lys Leu
168 20 25 30
169 aag aat agt gac tta agt ttt aat aat aaa gcg tta aaa gat cat aat 144
170 Lys Asn Ser Asp Leu Ser Phe Asn Asn Lys Ala Leu Lys Asp His Asn
171 35 40 45
172 gat gag tta act gaa gag ttg agt aat gct aaa gag aaa cta cgt gga 192
173 Asp Glu Leu Thr Glu Glu Leu Ser Asn Ala Lys Glu Lys Leu Arg Gly
174 50 55 60
175 tcc gcc gtg act agg ggt aca ata aat gac ccg caa aga gca aaa gaa 240
176 Ser Ala Val Thr Arg Gly Thr Ile Asn Asp Pro Gln Arg Ala Lys Glu
177 65 70 75 80
178 gct ctt gac aag tat gag cta gaa aac cat gac tta aaa act aag aat 288
179 Ala Leu Asp Lys Tyr Glu Leu Glu Asn His Asp Leu Lys Thr Lys Asn
180 85 90 95
181 gaa ggg tta aaa act gag aat gaa ggg tta aaa act gag aat gaa ggg 336
182 Glu Gly Leu Lys Thr Glu Asn Glu Gly Leu Lys Thr Glu Asn Glu Gly
183 100 105 110
184 tta aaa act gag aat gaa ggg tta aaa act gag gtc gac aga gtg ttt 384
185 Leu Lys Thr Glu Asn Glu Gly Leu Lys Thr Glu Val Asp Arg Val Phe
186 115 120 125
187 cct agg ggg acg gta gaa aac ccg gac aaa gca cga gaa ctt ctt aac 432
188 Pro Arg Gly Thr Val Glu Asn Pro Asp Lys Ala Arg Glu Leu Leu Asn
189 130 135 140
190 aag tat gac gta gag aac tct atg tta caa gct aat aat gac aag tta 480
191 Lys Tyr Asp Val Glu Asn Ser Met Leu Gln Ala Asn Asn Asp Lys Leu
192 145 150 155 160
193 cca tgg aga gtg cgt tat act agg cat acg cca gaa gat aag cta aaa 528
194 Pro Trp Arg Val Arg Tyr Thr Arg His Thr Pro Glu Asp Lys Leu Lys

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198          180          185          190
199    aat gag aag tta tct ctg cag aac ggt gat ggt aat cct agg gaa gtt 624
200    Asn Glu Lys Leu Ser Leu Gln Asn Gly Asp Gly Asn Pro Arg Glu Val
201          195          200          205
202    ata gaa gat ctt gca gca aac aat ccc gca ata caa aat ata cgt tta 672
203    Ile Glu Asp Leu Ala Ala Asn Asn Pro Ala Ile Gln Asn Ile Arg Leu
204          210          215          220
205    cgt cac gaa aac aag gac tta aaa gcg aga tta gag aat gca atg gaa 720
206    Arg His Glu Asn Lys Asp Leu Lys Ala Arg Leu Glu Asn Ala Met Glu
207    225          230          235          240
208    gtt gca gga aga gat ttt aag aga gct ggt acc ttg tta gat cag gtt 768
209    Val Ala Gly Arg Asp Phe Lys Arg Ala Gly Thr Leu Leu Asp Gln Val
210          245          250          255
211    aca caa tta tat act aaa cat aat agt aat tac caa caa tat aat gca 816
212    Thr Gln Leu Tyr Thr Lys His Asn Ser Asn Tyr Gln Gln Tyr Asn Ala
213          260          265          270
214    caa gct ggc aga ctt gac ctg aga caa aag gct gaa tat cta aaa ggc 864
215    Gln Ala Gly Arg Leu Asp Leu Arg Gln Lys Ala Glu Tyr Leu Lys Gly
216          275          280          285
217    ctt aat gat tgg gct gag agg ctg tta caa gag tta aat atc gat gtc 912
218    Leu Asn Asp Trp Ala Glu Arg Leu Leu Gln Glu Leu Asn Ile Asp Val
219          290          295          300
220    gcg act agg tct cag aca gat act ctg gaa aaa gta caa gaa cgt gct 960
221    Ala Thr Arg Ser Gln Thr Asp Thr Leu Glu Lys Val Gln Glu Arg Ala
222    305          310          315          320
223    gac aag ttt gag ata gaa aac aat acg tta aaa ctt aag aat agt gac 1008
224    Asp Lys Phe Glu Ile Glu Asn Asn Thr Leu Lys Leu Lys Asn Ser Asp
225          325          330          335
226    tta agt ttt aat aat aaa gcg tta aaa gat cat aat gat gag tta act 1056
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228          340          345          350
229    gaa gag ttg agt aat gct aaa gag aaa cta cgt aaa aat gat aaa tca 1104
230    Glu Glu Leu Ser Asn Ala Lys Glu Lys Leu Arg Lys Asn Asp Lys Ser
231          355          360          365
232    cta tct gaa aaa gct agt aaa att caa gaa tta gag gca cgt aag 1149
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242    fusion gene sequence constructed from
243    streptococcal type 24, 5, 6, 19, 1 and 3 M protein
244    DNAs

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VERIFICATION SUMMARY  
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Original Text

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